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## ALIGNMENTS

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REFERENCE AUTHORS TITLE JOURNAL COMMENT SOURCE ORGANISM RESULT 1 BM451059 LOCUS KEYWORDS VERSION ACCESSION DEFINITION Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
CDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information car
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAMI2121 row: d column: 08 Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 889 bp mRNA linear E AGENCOURT\_6392456 NIH\_MGC\_67 Homo sapiens cDNA clone 5', mRNA sequence. BM451059 1 (bases 1 to 889)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999) Homo sapiens (human) BM451059.1 GI:18500099 quality sequence stop: 665 EST 05-FEB-2002 e IMAGE:5494999 can þ

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Description

SUMMARIES

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80.6 77.1 75.1 74.4

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BM451059 CB852030 BM051236 CA777547

BM451059 AGENCOURT CB852030 UI-CF-FN0 BM051236 603634191 CA777547 ip21g07.x

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/tissue type="retinoblastoma"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_67"
/note="Organ: eye; Vector: pcMV-SPORT6; Site_1: Not1;
Site_2: Sall; Cloned unidirectionally. Primer: Oligo
Average insert size 1.75 kb. Library constructed by
Technologies."
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771 bp mRNA linear EST 22-Ai UI-CF-FNO-aes-j-02-0-UI.sl UI-CF-FNO Homo sapiens cDNA clone UI-CF-FNO-aes-j-02-0-UI 3', mRNA sequence. CB852030 CB852030.1 GI:30046884
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Tissue Procurement: Dr. M. J. Welsh, University of Iowa
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
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/tissue type="Human Lung Epithelial cells"
/lab host="DH108 (Life Technologies) (Tl phage resistant)
/lab host="DH108 (Life Technologies) (Tl phage resistant)
/clone lib="UI-CF-FN0"
/note="Organ: Lung; Vector: pT773-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-CF-FN0 is a subtracted cDNA library derived from two
normalized Human lung epithelial cell libraries (EN1 and
DU1) The library was subtracted according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
Bonaldo, For additional information, contact:
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                                                                         found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM1886 row: e column: 23
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1 (bases 1 to 814)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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BM051236.1
                                                                                                                                                                                                                                                                                                                                                   603634191F1 NIH_MGC_43 Homo sapiens cDNA clone IMAGE:5424766 5',
                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                           Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                              Homo sapiens (human)
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/clone_lib="NIH MGC 43"
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2: RORI; cDNA made by Oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5; adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library. | "
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/tissue_type="normal pigmented retinal epithelium"
/lab_host="DH10B (phage_resistant)"
/lama_lih="NIH_MGC_43"
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Library was constructed by Dr. Hiroshi Inoue DNA sequencing
Washington University Genome Sequencing Center For informati
Obtaining a clone please contact: Dr. Hiroshi Inoue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Douglas Melton, Klaus H. Kaestner, & Hiroshi Inoue
Endocrine Pancreas Consortium
Harvard University, Howard Hughes Medical Institute
Dept of Molecular and Cellular Biology, 7 Divinity Ave, Cambridge,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Melton,D., Brown,J., Kenty,G., Permutt,A., Lee,C., Kaestner,K., Lemishka,I., Scearce,M., Brestelli,J., Gradwohl,G., Clifton,S., Hillier,L., Marra,M., Pape,D., Wyle,T., Martin,J., Blistain,A., Schmitt,A., Theising,B., Ritter,E., Ronko,I., Bennett,J., Cardenas,M., Gibbons,M., McCann,R., Cole,R., Tsagareishvili,R., Williams,T., Jackson,Y. and Bowers,Y. Endocrine Pancreas Consortium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (hinoue@im.wustl.edu)
Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 617-495-1812
Fax: 617-495-8557
                                                                                                                                                                                                                                                                                                                                      Similarity
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756 bp mRNA linear EST 03-DEC-2002 ip21g07.x1 HR85 islet Homo sapiens cDNA clone IMAGE:6217933 3' similar to TR:085769 085769 HYPOTHETICAL 24.8 KD PROTEIN. ;, mRNA
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                                                                                                                                                                                                     CTTCCCCCGAGGACAGCCGCCTGTGGCAGTATCTTTTGAGCCGCTCCATGCGGGAGCAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Size-selected on agarose gel. Average insert size -1kb. 5' XhoI site was destroyed after directional cloming. Amplified once Contact information: Hiroshi Inoue, MD, Metabolism Div. (Alan Permutt Lab), Washington University School of Medicine, Box 8127, 660 South Euclid Ave., St. Louis, MO 63110, E-mail: hinoue@imgate.wustl.edu, Tel: 314-362-1916, Fax: 314-747-2692."
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                                                                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                                                                                                                                                                                     1 (bases 1 to 1006)
NIH-MGC http://mgc.nci.nih.gov/.
Mational Institutes of Health, Mammalian Gene
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                         guality sequence stop: 745.
                                                                                                                                                                                                                                                                                                                                                                                                             sapiens
/organism="Homo sapiens"
/mol_type="mRNA"
                                                           Location/Qualifiers
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cDNA clone IMAGE:4297911
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                                                                               CATCCGGCGGGACGTCAGGGTCTACATCAGCCTCCT-GCCCCTGGGCGATGGACTCACC-
                                            CATCCGGCGGGACGGCAGGGTCTACATCAGGCTTCTGGCCCCTGGGCGATGAGATCACCC
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/clone_lib="NH10B (phage-resistant)"
/clone_lib="NH1MGC_48"
/note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into BccRI/KhoI sites using the following 5' adaptor: GGCAGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis with (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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/clone="IMAGE:4297911"
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90.9%;
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Pred. No. 1.4e-79;
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TITLE
JOURNAL
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                                                                                                                                                N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.B. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health, I
Unpublished (1999)
                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Plate: LLCM1182 row: h column:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            602145812F1 NIH_MGC_48 Homo sapiens cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BF664198.1 GI:11938093
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                                                               CCGCACTGGGCGCCTTCGCCACTGGCCTCTTCCTGGGGAGGCGGTGCCCCCATGGC
                                                                                         CCGCACTGGGCCGCCTTCGCCACTGGCCTTCCTTCGGGGAGGCGGTGCCCCCCATGGC
                                                                                                                                            GCGCCATGACCCAGCCGGTGCCCCGGCTCTCCGTGCCCGCCGCGCTGGCCCTGGGCTCAG 138
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                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                         /clone="IMAGE:4309298"
/clone="IMAGE:4309298"
/tissue_type="primary B-cells from tonsils (cell line)"
/tissue_type="primary B-cells from tonsils (cell line)"
/lab_host="pH10B (phage-resistant)"
/clone_lib="NIH_MGC 48"
/clone_lib="NIH_MGC 48"
/note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
Site_2: ECRI; cDNA made by oligo-dT priming.
Directionally cloned into EcRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:4309298"
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97.4%;
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                                                                                                                                                                                                                                             Score 716; DB 10;
Pred. No. 5.9e-79;
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/lab host="DH108 (Life Technologies) (T1 phage resistant)"
/clone_lib="UI-CF-EN1"
/clone_lib="UI-CF-EN1"
/note="Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-CF-EN1 is a normalized cDNA library containing the following tissue(s): Primary Lung Cystic Fibrosis
Epithelial Cells. The library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into pT7T3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library lag sequence that is located between the Not I site and the (dT)18 tail. The sequence tag for this library is CTGCTCAGGT.

TAG TISSUS-Human Lung Epithelial Cell Lines untreated LPS for the class of the contains a library tag sequence that is the contains a library the contains and the contains a library tag sequence that is library tag sequence that is sequence tag for this library is CTGCTCAGGT.
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="UI-CF-ENI-acy-e-23-0-UI"
/tissue_type="Primary Lung Cystic Fibrosis Epithelial
Cells"
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                                                                                                                                                                                                                                                                                                                                                                                                                          Email: paul-mccray@uiowa.edu
Tissue Procurement: Dr. M. J. Welsh, University of Iowa
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
cDNA Sequencing by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2024 University of Iowa Med Labs, Iowa City, Tel: 319 356 4866 Fax: 319 356 7171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UI-CF-FNO-afl-p-08-0-UI.S1 UI-CF-FNO Homo sapiens cDNA clone UI-CF-FNO-afl-p-08-0-UI 3', mRNA sequence.
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Bonaldo, M.F., Lenno
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                    bento-soares@uiowa.edu
TAG_SEQ=None found"
                                                                                                                                                                            /tissue_type="Human Lung Epithelial cells"
/lab_host="DH10B (Life Technologies) (T1 p:
/clone_lib="UI-CF-FN0"
                                                                                                                                                                                                                                    clone="UI-CF-FN0-afl-p-08-0-UI"
                                                                                                                                                                                                                                                         organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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RESULT 9 BE796570 LOCUS

DEFINITION

601592247F1 NIH\_MGC\_7 Homo

772 bp sapiens

mRNA cDNA

clone

linear BST 20-9 one IMAGE:3946546

20-SEP-2000

ACCESSION VERSION

KEYWORDS

EST

BE796570.1 mRNA sequence. BE796570

GI:10217768

SOURCE ORGANISM

Homo sapiens Homo sapiens (human)

REFERENCE

Bukaryota; Metazoa; Chordata; C Mammalia; Eutheria; Primates; C 1 (bases 1 to 772) NIH-MGC http://mgc.nci.nih.gov/

Primates; Catarrhini; Hominidae;

Craniata; Vertebrata; Euteleostomi;

AUTHORS

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                 TCTAGGGCTGGCCCCTAGTGAGTGGGCTCGAG 899
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                                                                                                      TGCGACCCGGAGGCATCCTCGCCGTCCTCAGAGTCCTGTGGCGCGGGAAGGTGCTGCAAC
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                                                   ACGTCAGGGTCTACATCAGCCTCCTGCCCCTGGGCGATGGACTCACCTTGG-CTTCAAGA
                                                                             ACGTCAGGGTCTACATCAGCCTCCTGCCCCTGGGCGATGGACTCACCTTGGCCTTCAAGA
                                                                                                                                 CCGCCTTGGAGACCCTGGACGAGCTGCTGGCGGCGGCGAGGCCGGCACCTTCGACGTGG
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Tissue Procurement: DCTD/DTP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)
GGAGCTGGGACGGCCCCTGTGGAGGCAGGCAGGAGCACAAGATCGACCTCCGGCT
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/clone lib="NIH MGC 7"
/clone Torgan: lung; Vector: pOTB7; Site 1: XhoI; Site 2:
/note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site 2:
/note="Organ: lung; Vector: poTB7; Site 1: XhoI; Site 2:
/note="Organ: lung; Vector: poTB7; Site 1: XhoI; Site Site 3: XhoI; Site Site 3: XhoI; Site Site 3: XhoI; Site Site 3: XhoI; Site Site Site 4: XhoI; Site Site Site 4: XhoI; Site Site Site 4: XhoI; Site Site Site 3: XhoI; Site Site Site 3: XhoI; Site Site Site 3: XhoI; Site Site 3: XhoI; Site Site 3: XhoI; Site Site 3: XhoI; Site 3: XhoI
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/lab_host="DH10B (phage-resistant)"
/-1--- 14h-"NTH MGC 7"
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/cell_line="MGC3"
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Pred. No. 4.7e-75;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1181 row: 1 column: 03
High quality sequence stop: 761.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
CDNA Library Preparation: Ling Hong/Rubin Laborator
CDNA Library Arrayed by: The I.M.A.G. Consortium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 779)
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                                                                                                                                                                                                                                                                                     /tissue_type="primary B-cells from tonsils (cell line)"
/lab_hoSt="DH10B (phage-resistant)"
/clone lib="NNIH_MGC_48"
/clone lib="NNIH_MGC_48"
/note="Togan: B-cells; Vector: pOTB7; Site_1: Xho1;
/note="Togan: B-cells; Vector: pOTB7; Site_1: Xho1;
/note="Togan: B-cells; Vector: pOTB7; Site_sites using the Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Lir Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit
                                                                                                                                                                                                                                                California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM2613 row: e column: 24
                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D
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BQ931589.1 GI:22346620
                                                             cDNA Library Preparation: Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
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/note="Organ: lung; Vector: pOTB7; Site_1: EcoRI; Site_2: /note="Organ: lung; Vector: potentially cloned into EcoRI/XhoI sites using the following 5: adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
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cDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
lateral Column: 17
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NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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Tissue Procurement: Life Technologies, Inc.
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /CLUME 1119 ALM PAGE 112 ALM PAGE 112 ALM PAGE 12 ALM PAGE 12 PCRT (destroyed); RNA source anonymous pool of 24 week female lung, 16 week female spleen, and 20-22 week male spleens. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.4 kb, insert size range 1-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Grubber (Invitrogen). Research Genetics tracking code 026. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'mol_type="mRNA"
'db_xref="taxon:9606"
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Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                                                                             52 Eoeun-dong Yuseong-gu,
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 625)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and Kim,Y.S.

Kim,Y.S.

Z1C Frontier Korean EST Project 2001
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/note="Organ: Stomach; Vector: pTZ18RP1; Site_1: EcoRI;
Site_2: NotI; The poly (A)+ RNA was dephosphorylated with
                                 /lab_host="Top10F'"
/clone_lib="S14K402"
/note="Organ: Stomacl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         63.2%; Score 625; DB 12; 100.0%; Pred. No. 1e-67; ive 0; Mismatches 0;
                                                                                                                                                                                                     mRNA
                                                                                                                                                         cDNA clone
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E:4555609 5',
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ORGANISM
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Matches
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Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1256 row: o column: 02
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D
CGCAGCCCCCGGAGCTGGGACGCCCCTGTGGAGGCAGAGGCCGAGGCGCAGAGATCG
                                                                                                                                      TGGCGCGGCTCATCCAGGCCAAGAAGGCGCTGGACCTGGGCACCTTCACGGGCTACTCCG
                                                                                                                                                                                                                                                                                                                                                                                            GCTCAGCCGCACTGGGCGCCGCCTTCGCCACTGGCCTCTTCCTGGGGAGGCGGTGCCCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.B. Consortium (
                                                                                  CCCTGGCCCTGGCCCTGGCGCTGCCGCGGACGGCGTGGTGACCTGCGAGGTGGACG
                                                                                                               TGGCGCGGCTCATCCAGGCCAAGAAGGCGCTGGACCTGGGCACCTTCACGGGCTACTCCG
                                                                                                                                                                             TGGAGCAGCCGCAGGGGATTCTATGATGACCTGCGAGCAGGCCCAGCTCTTGGCCAACC
                                                                                                                                                                                                             TGGAGCAGCCGCAGGGGATTCTATGATGACCTGCGAGCAGGCCCCAGCTCTTGGCCAACC
                                                                                                                                                                                                                                             ATCTTCTGAGCCGCTCCATGCGGAGCACCCGGCGCTGCGAAGCCTGAGGCTGCTGACCC
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism≃"Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 621; DB 12;
Pred. No. 2.8e-67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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711 AACGCAT-CGGCGGGACGTCAGGGTCTACATCAGCCTCCTGCCCTGGCG
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                                      The FANTOM Consortium and the RIKEN Genome Exploration Research Group Phase I & II Team.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y. Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genes Genome Res. 10 (10), 1617-1630 (2000)
         Group Phase I \& II Team. Analysis of the mouse transcriptome based on functional annotation
                                                                                                                                                                                              FANTOM Consortium.
                                                                                                                                                                                                                    The RIKEN Genome Exploration Research Group Phase II Team and
                                                                                                                                                                     Functional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AKO54334 20-33 bp mRNA linear HTC 20-SEP-20 MUS musculus 2 days pregnant adult female ovary CDNA, RIKEN full-Length enriched library, clone:E330016LD2 product.weakly similar to PUTATIVE O-METHYLTRANSFERASE (FRAGMENT) [Rhodothermus
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AK054334
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                                                                                                                                                  annotation of a full-length mouse cDNA collection
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                                                                                                                                     685-690
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                                                                                                                                     (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Rike Genomic Sciences Center and Genome Science Laboratory in RIKEN.

Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Exploration Research Group, RIKEN Genomic Sciences Center (GSC) RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama Gado-ouds, Japan (E-mail:genome-res@gsc.riken.go.jp, URL:http://genome.gsc.riken.go.jp/, Tel:81-45-503-9222, Fax.81-45-503-9216)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      URL:http://genome.gsc.riken.go.jp/
URL:http://fantom.gsc.riken.go.jp/.
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Please visit our web site for further details.
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Nature 420, 563-573 (2002)
6 (bases 1 to 2243)
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/db_xref="01:26344157"
/db_xref="01:26344157"
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/tzenftion="mappyprisipaalalagsaalgaafargliligkrwppwgsrrqe
/tzenftion="mappyprisipaalalagsaalagsaalgaafargliligkrwppwgsrrqe
RLLPPEDNPLWQYLLSRSWREHPALRSLRLTLTLEQPQGDSMMTCEQAQLLANLARLIK
AKKALDLGTFTGYSALALALALPEAGRVVTCEVDAEPPKLGRPMWKQAEVEQKIDLRL
QPALQTLDELLAAGEAGTFDIAVVDADKENCTAYYERCLQLLRPGGVLAVLRVLWRGE
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weakly similar to PUTATIVE O-METHYLTRANSFERASE
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                                                                                                                                                                                                                                                                                                                                                                          /note="putative"
                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="putative"
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40. .828
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="ovary"
/clone_lib="RIKEN full-length enriched mouse cDNA library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /codon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                [Rhodothermus marinus]
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          Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning
Meth. Enzymol. 303, 19-44 (1999)
                                                                                                                                                                                                                          Mus
Mus
                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus 10 day old male pancreas cDNA, RIKEN full-length enriched library, clone:1810030M08 product:weakly similar to PUTATIVE O-METHYLTRANSFERASE (FRAGMENT) [Rhodothermus marinus],
                                                                                                                                                                                   Eukaryota;
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Group Phase I & II Team.

Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs

AL Nature 420, 563-573 (2002)

CB 6 (bases 1 to 919)

RS Adachi, J., Alzawa, K., Akahira, S., Akimura, T., Arai, A., Aono, H., Arakawa, T., Bono, H., Carninci, P., Fukuda, S., Fukunishi, Y., Furuno, M., Hanagaki, T., Hara, A., Hayatsu, N., Hiramoto, K., Hiraoka, T., Kato, H., Ishi, Y., Itoh, M., Izawa, M., Kasukawa, T., Kato, H., Kawai, J., Kojima, Y., Konno, H., Kouda, M., Koya, S., Kurihara, C., Matsuyama, T., Miyazaki, A., Nishi, K., Nomura, K., Nunazaki, R., Ohno, M., Okazaki, Y., Okido, T., Owa, C., Salto, H., Salto, R., Sakai, C., Sakai, X., Saho, H., Sasaki, D., Salto, R., Sahinagawa, A., Shiraki, T., Sogabe, Y., Suzuki, H., Tagami, M., Tagawa, A., Takahashi, F., Tanaka, T., Yoshina, Y., Toya, T., Yamamura, T., Yasunishi, A., Yoshida, K., Yoshinassion
                                                                                                  Submitted (10-UUL-2000) Yoshihide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Please visit our web site (http://genome.gsc.riken.go.jp/) further details.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KRIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokanagawa 230-0045, Japan (E-mail:genome-res@gsc.riken.go.jp, URL:http://genome-gsc.riken.go.jp/, Tel:81-45-503-9222, Fax:81-45-503-9216)
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/dev_stage="10 day old"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8.8
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/strain="C57BL/60"
/db_xref="FANTOM_DB:1810030M08"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="1810030M08"
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80.8%;
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Pred. No. 1.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches 169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.2e-65;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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           902
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ORGANISM
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AW157329/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACCESSION
VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DEFINITION
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                                                                            Matches
                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE
JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AUTHORS
                                                                          604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jos Krizman, D., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R. WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AW157329
606 bp mRNA linear EST 04-NOV-1999
au94f01.x1 Schneider fetal brain 00004 Homo sapiens cDNA clone
IMAGE:2783929 3' similar to TR:085769 085769 HYPOTHETICAL 24.8 KD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Washington University School
4444 Forest Park Parkway, Box
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HOMO sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AW157329.1 GI:6228730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROTEIN. ;, mRNA sequence.
AW157329
                            GAGCAGGCCCAGCTCTTGGCCAACCTGGCGCGCTCATCCAGGCCAAGAAGAAGGCGCTGGAC
GACCAGGCCCAGCTCTTGGCCAACCTGGCGCGCTCATCCAGGCCAAGAAGGCGCGCTGGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GATGGCCTCTCGTTGGCCTTTAAGATCTAGGGTTAACACGAAGCTTAGGGCCT---GGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTGAGAACCTAAGACCCCAGCCAGTGACCTGACTTTTAAACCTGACAATAAAG 906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGGTTGCCTGGGAACCCCCAGGAATTGACCCCTGAGTTTTAAATTCGAAAATAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 606)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer: -40UP from Gibco
h quality sequence stop: 443.
Location/Qualifiers
                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              314 286 1800
314 286 1810
                                                                                                                                       /note="Organ: brain; Vector: pBluescript SK (Stratagene);
Site_1: SstI; Site_2: XhoI; Double-stranded cDNA was
prepared from human fetal brain tissue. 5' and 3'
                                                                                                                                                                                                                                                                                                        adaptors were used in cloning as follows: 5' adaptor
                                                                                                                                                                                                                                                                                                                                                                                                                       clone
                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="5 months post-conception"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue_type="frontal lobe"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              clone="IMAGE:2783929"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"
|mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Metazoa;
                                                                                           61.0%;
99.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                   _lib="Schneider fetal brain 00004"
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Primates;
                                                                      0;
                                                                      Score 603.4;
Pred. No. 4.5e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Box
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x 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                         4.5e-65;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dubuque, T., Geisel, G., Jost,
                                                                                                        DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Louis,
                                                                      Indels
                                                                                                      Length
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REFERENCE
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BY707694
LOCUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM
Schazaki, Y., Furunc, M., Kasukawa, T., Adachi, J., Bono, H., Kondo, S., Nikaido, I., Osato, N., Saito, R., Suzuki, H., Yamanaka, I., Kiyosawa, H., Yagi, K., Tomaru, Y., Hasegawa, Y., Nogami, A., Schonbach, C., Gojobori, T., Baldarelli, R., Hill, D.P., Bult, C., Hume, D.A., Quackenbush, J., Schriml, L.M., Kanapin, A., Matsuda, H., Batdavo, S., Beisel, K.W., Blake, J.A., Bradt, D., Brusic, V., Chothia, C., Corbani, L.E., Cousins, S., Dalla, E., Dragani, T.A., Fletcher, C.F., Forrest, A., Frazer, K.S., Gaasterland, T., Garimond, S., Hirokawa, N., Jackson, I.J., Jarvis, E.D., Kanai, A., Kawaji, H., Kawasawa, Y., Kedzierski, R.M., King, B.L., Konagaya, A., Kurochkin, I.V., Lee, Y., Lenhard, B., Lyons, P.A., Maglott, D.R., Maltais, L., Marchionni, L., McKenzie, L., Miki, H., Nagashima, T., Numata, K., Okido, T., Pavan, W.J., Pertea, G., Pesole, G., Petrovsky, N., Pillai, R., Pontius, J. U., Qi, D., Ramachandran, S., Ravasi, T., Reed, J.C., Reed, D.J., Raid, J., Ring, B.Z., Ringwald, M., Sandelin, A., Schneider, C., Sepple, C.A., Setou, M., Shimada, K., Sultana, R., Takenaka, Y., Taylor, M.S., Teasdale, R.D., Tomita, M., Verrardo, R., Wagner, L., Wahlestedt, C., Wang, Y., Watanabe, Y., Watanabe, Y., Watanabe, Y., Wahlestedt, C., Wang, Y., Watanabe, Y., Watanaba, Y., Watanabe, Y., Watanaba, Y., Watanabe, Y., Watanabe,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
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                                                                                                                                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Computational Analysis of Full-Length Mouse cDNAs Compared with Human Genome Sequences Mamm. Genome. 12, 673-677 (2001)

Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genes. Genome Res. 10 (10), 1617-1630 (2000)

RIKEN integrated sequence analysis (RISA) system-384-format sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 1757-1771 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp/
Adachi_J., Aizawa,K., Akimura,T., Arakawa,T., Carninci,P.,
Fukuda,S., Hashizume,W., Hayashida,K., Hirozane,T., Hori,F.,
Fukuda,S., Hashizume,W., Hayashida,K., Hirozane,T., Kojima,Y.,
Imotani,K., Ishii,Y., Itoh,M., Kagawa,I., Kawai,J., Kojima,Y.,
Kondo,S., Konno,H., Koya,S., Myazaki,A., Murata,M., Nakamura,M.,
Nomura,K., Numazaki,R., Ohno,M., Ohsato,N., Saito,R., Sakazume,N.,
Takeda,Y., Waki,K., Watahiki,A., Muramatsu,M. and Hayashizaki,Y.
                                                                                                                                                                                                                                                         Similarity
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The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suebiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
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                                                                       CCGCCGCCTCCTGCCCGCGCATGACCCAGCCGGTGCCCCGGCTCTCCGTGCCCGCCGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Please visit our web site (http://genome.gsc.riken.go.jp) for
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /tissue_type="pancreas"
/dev_stage="10 day old"
/clone_lib="RIKEN full-length enriched, 10 day old
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="male"
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/db_xref="taxon:10090"
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JOURNAL COMMENT
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AUTHORS
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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//clone_lib="Soares_NFL_T_GBC_S1"
//note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with
/note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with
a modified polylinker; Site 1: Not I; Site 2: Eco RI;
Equal amounts of plasmid DNA from three normalized
libraries (fetal lung NbHL19W, testis NHT, and B-cell
NCI_CGAP_GCB1) were mixed, and ss circles were made in
vitro. Following HAP purification, this DNA was used as
tracer in a subtractive hybridization reaction. The driver
was PCR-amplified cDNAs from pools of 5,000 clones made
from the same 3 libraries. The pools consisted of
I.M.A.G.B. clones 297480-302087, 682632-687239,
726408-728711, and 729096-731399. Subtraction by Bento
Soares and M. Fatima Bonaldo. "
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/db_xref="taxon:9606"
/clone="IMAGE:2330365"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 1122)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999)
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                                                                                               AACCTG----GCGCGGCTCATCCAGGCCAAGAAGGCGCTGGAACCTGGGCACCTTCACGGGC
                                                                                                                                                                         CCCTGGAGCAGCCGCAGGGGGATTCTA-TGATGACCTGCGAGCAGGCCCAGCTCTTGGCC
                                                                                                                                                                                                                                                        GTATCTTCTGAGCCGCTCCATGCGGGAGCCCCGG-CGCTGCGAAGCCTGAGGCTGCTGA
                                                                                                                                                                                                                                                                                                                                                                            GCTCAGCCGCTACTGGGCGCGCGTTCGCCACTGGCCTCTTCCTGGGGAGGCGGTGCCCC
                                                              AACCTGTGCGCGGCTCTATCCAGGCCAATGAAGGCGCTGGACCTGGGCACCTTCACGGGC
                                                                                                                                          CCCTGGAGCAGCCGCAGGGGATTCTAGTGATGACCTGCGAGCAGGCCCCAGCTCTTGGAC 301
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/clone="IMAGE:455585"
/tissue_type="leiomyosarcoma cell line"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH MGC 46"
/clone_lib="NIH MGC 46"
/note="Organ: uterus; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="CONA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GGCACGAG(G). Size-selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-cDNA synthesis kit (Stratagene) and Superscript
IRT (Life Technologies). Note: this is a NIH_MGC
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/db_xref="taxon:9606"
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87.2%;
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RESULT 21
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                                                                                                                                                                                                Genome Research Center
Genome Research Institute of Bioscience & Biotechnology
Korea Research Institute of Bioscience & Biotechnology
Formun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                             Email:
Plate:
                                                                                                                                                                                                                                                                                   Kim,Y.S.
21C Frontier Korean EST Project
                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (Dases 1 to 580)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S.
                                                                                                                                                                            52 Eoeun-dong Yuseong-gu,
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                   Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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K-EST0052976 S14K402 Homo
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Location/Qualifiers
                                                                                                                                               yongsung@mail.kribb.re.kr
22 row: C column: 12
/organism="Homo sapiens"
/mol type="mRNA"
/db xref="taxon:9606"
/clone="S14K402-22-C12"
/cell_line="K402"
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sapiens
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S. and
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RESULT 22
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5', mRNA sequence.
BQ918468
BQ918468 1 GI:22333166
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/clone lib="914K402"
/clone lib="914K402"
/note="Torgan: Stomach; Vector: pTZ18Rp1; Site_1: EcoRI;
Site_2: NotI; The poly (A) + RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand obtained cDNA vectors were used for transformation of competent cells E. coli TOplOF; by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
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Pred. No. 3.5e-62;
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                                                                           sapiens cDNA clone IMAGE: 6429860
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KEYWORDS

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AUTHORS
TITLE
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Best Local Similarity
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718 GAGTCCTGTGGCGCGGGAAGGTGCTGCAACCTCCGAAAGGGGACGTGGC-GGCCGAGTGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM2618 row: 1 column: 21
High quality sequence start: 12
High quality sequence stop: 509.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteresto Mammalla; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 998)

NIH-MGC http://mgc.nci.nih.gov/.

Mational Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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Tissue Procurement: DCTD/DTP/Gazdar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                         CCGCCTACTACGAGCGCTGCCTGCAGCTGCTGCGACCCGGGAGCATCCTCGCCGTCCTCA
                                                                                                           CCGCCTACTACGAGCGCTGCCTGCAGCTGCTGCGACCCGGAGGCATCCTCGCCGCCGTCCTCA
                                                                                                                                                                      CGGCGGGCGAGGCCGTTCGACGTGGCCGTGGTGGATGCGGACAAAGAGAACTGCT
                                                                                                                                                                                                                CGGCGGGCGAGGCACCTTCGACGTGGCCGTGGTGGATGCGGACAAGGAGAACTGCT
                                                                                                                                                                                                                                                                                                             CGGAGCACAAGATCGACCTCCGGCTGAAGCCCGCCTTGGAGACCCCTGGACGACGAGCTGCTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="large cell carcinoma"
/lab_host="PH10B (phage-resistant)"
/clone_lib="NIH_MGC_18"
/clone_lib="NIH_MGC_18"
/note="Organ: lung; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into BcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other_ESTs: au59e03.x1
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AI929359

BY MRNA linear EST 23-AUG-
au59e03.yl Schneider fetal brain 00004 Homo sapiens cDNA clone
IMAGE:2519068 5' similar to TR:085769 085769 HYPOTHETICAL 24.8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Wilson RK
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h quality sequence stop: 388.
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314 286 1810
and 3' adaptor sequence:
5'-GAGAGAGAGACCGAGTTTTTTTTTTTTTTTT-3'. The library was
size-selected for >0.5 kb inserts and has an average
insert size estimated at 1.2 kb. This library was
constructed using the CAP-trapper method for full-length
                                                                                                                                                                          /note=Torgan: brain; Vector: pBluescript SK (Stratagene); Site 1: SstI; Site 2: XhoI; Double-stranded cDNA was prepared from human fetal brain tissue. 5' and 3'
                                                                                                                  adaptors were used in cloning as
                                                                                                                                                                                                                                                /clone_lib="Schneider fetal brain 00004"
                                                                                                                                                                                                                                                                      /dev_stage="5 months post-conception"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                 tissue_type="frontal lobe"
                                                                                                                                                                                                                                                                                                                                         sex="male"
                                                                                                                                                                                                                                                                                                                                                             clone="IMAGE:2519068"
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x 8501, St. Louis,
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                                                                 TGGACGAGCTGCTGGCGGCGGCGGCAGGCCGCACCTTCGAC 623
TGGACGAGCTGCTGGCGGCGCGCGAGGCCGGCACCTTCGAC 581
                                                                                                                                         GGAGGCAGGCCGAGCACAAGATCGACCTCCGGCTGAAGCCCGNCTTGGAGACCC
                                                                                                                                                                                                              GGAGGCAGGCCGAGGCGAGACCAAGATCGACCTCCGGCTGAAGCCCGCCTTGGAGACCC
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Pred. No. 5.4e-62;
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REFERENCE AUTHORS SOURCE ORGANISM ACCESSION VERSION KEYWORDS Mammalia; Butheria; Primates; Catarrhini;
1 (bases 1 to 691)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian G
Unpublished (1999) Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Homo sapiens mRNA sequence. BE279279 BE279279.1 GI:9154269 (human) Hominidae; Collection (MGC) Euteleostomi;

COMMENT

Email: cgapbs-r@mail.nih.gov Tissue Procurement: ATCC

cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (1) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information

(LINL)

can

Contact: Robert Strausberg, Ph.D

RESULT 24 BE279279

DEFINITION

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mRNA cDNA clone

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EST 13-JUL-2000

IMAGE:3504600

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                                                                                                                                                                                                        GAGCTGGGACGGCCCTGTGGAGGCAGGCCGAGGCGGAGCACAAGATCGACCTCCGGCTG
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CTGCTGCGACCCGGAG
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                        CTGCTGCGACCCGGAG
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Directionally cloned into EcoRI/KhoI sites using the following 5: adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="choriocarcinoma"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NHH_MGC_21"
/note="Organ: NHACENTA Vector: pOTB7; Site_1: XhoI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:9606"
/clone="IMAGE:3504600"
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Pred. No. 5.2e-62;
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plate: NDAM0034 row: o column: 18
High quality sequence stop: 622.
Location/Qualifiers
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CDNA Library Preparation: Invitrogen Corp
CDNA Library Arrayed by: The I.N.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
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Unpublished (1999)
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NIH-MGC http://mgc.nci.nih.gov/.
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                                    ACCTTCACGGGCTACTCCGCCCTGGCCCTGGCCCTGGCCCCGCGGACGGGCGCGTG
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ACTTTCACGGGCTACTCGGCCCTGGCCCTAGCCTTGGCGCTTCCGGAGGCTGGCCGCGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="DH10B (phage-resistant)"
/clone lib="NIH MGC 135"
/clone lib="NIH MGC 135"
/note="Vector: pcMV5port6.1; Site 1: EcoRV; Site 2: NotI;
/note="Vector: pcMV5port6.1; Site 2: NotI;
/notInent: NotInent: NotIn
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/db_xref="taxon:10090"
/clone="IMAGE:30136001"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Mus"
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Rodentia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 574.6; DB 14;
Pred. No. 1.4e-61;
0; Mismatches 167;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalian Gene Collection (MGC)
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AW157252/c
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                                                                                                                            Seq primer: -40UP from Gibco
High quality sequence stop: 4
                                                                                                                                                            This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                               Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1 Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 582)

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R. WashU-KCI human EST Project
                                                                                                                                                                                                              Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AW157252 582 bp mRNA linear EST 04-NOV-1999 au93e01.x1 Schneider fetal brain 00004 Homo sapiens cDNA clone IMAGE:2783832 3' similar to TR:085769 085769 HYPOTHETICAL 24.8 KD
                                                                                                                                                                                                                                                                                                                                         Unpublished (1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                  Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AW157252.1 GI:6228653
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                           /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
            /clone="IMAGE:2783832"
                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                     Louis,
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CB181160 886 bp mRNA 1 AGENCOURT 11381817 NIH MGC 164 Mus musculus IMAGE:30240746 5', mRNA sequence.
CB181160 CB181160.1 GI:28177885
                                                                                                                                                                                                                                                                                        GGGAACCCCAGGAATTGACCCTGAGTTTTAAATTCGAAAATA 953
                                                                                                                                                                                                                                                       GAACGCATCCGGCGGGACGTCAGGGTCTACATCAGCCTCCTGCCCCTGGGCGATGGACTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone lib="Schneider fetal brain 00004"
/note="Organ: brain; Vector: pBluescript SK (Stratagene);
/site_1: SstI; Site_2: XhoI; Double-stranded cDNA was
prepared from human fetal brain tissue. 5' and 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptors were used in cloning as follows: 5' adaptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /sex="male"
/tissue_type="frontal lobe"
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/lab_host="DH10B"
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98.1%;
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Pred. No. 2.9e-60;
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National Institutes of Health, N
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 611
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                       found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
| Plate: LLCM2695 row: e column: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Michael Brownstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGENCOURT 10119399 NIH MGC_144 Mus musculus cDNA clone IMAGE:6535487 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria;
1 (bases 1 to 977)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAACCTGACGAACGCATCCTGAGGGACGCCAGGGTCTACATCAGCCTCTGCCCC
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         /lab_host="DH10B (T1-phage-resistant)"
/clone_lib="NHH_MGC 144"
/clone_lib="NHH_MGC 144"
/clone_lib="NHH_MGC 144"
/note="Organ: Brain; Vector: pDNR-LHB; Site_l: Sfil
/ggcgattatggcc); Site_2: Sfil (ggcgcatcggcc); cDNA made
by oligo-dT priming and directionally cloned. 5' and 3'
adaptors were used in cloning as follows;
5',ARGCAGTGGTATCAACGCAGAGTGGCATTACGGCCGGG-3' and
5',ARTCTAAGGCCGACGACGACTG-dT(30)NN-3'. Full-length
enriched library was constructed using the Clontech
Creator SMART kit and size-selected to contain the 0.2-0.5
kb size fraction (other fractions present in NIH_MGC 143)
Library created in the laboratory of M. Brownstein (NIMH,
NIH). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                            /mol_type="mRNA"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                      organism="Mus musculus"
                                                                                                                                                                                                                                                                                                   clone="IMAGE: 6535487"
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Query Match

Best Local Similarity

56.3**%;** 

Score 557.2; DB 1 Pred. No. 1.9e-59;

DB 13;

Length 977;

Matches

679;

Conservative

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899
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                                                                                   578 bp mRNA linear EST 08-MAR-2000 wq66g10.x1 NCI CGAP GC6 Homo sapiens CDNA clone IMAGE:2476290 3' similar to TR:\overline{O85769} O85769 HYPOTHETICAL 24.8 KD PROTEIN.;, mRNA
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                      AW003514.1
EST.
                                                          sequence.
AW003514
   Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGGTGGCCTCCATGGGGGTCCAGGCGGCAAGAGCGCCTGCTGCCACCTGAGGACAATCCC
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DNA Sequencing by: Washington University Genome Sequencing C

Clone distribution: NCI-CGAP clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 799 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tissue Procurement: Christopher A. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia; Eutheria;
1 (bases 1 to 578)
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Mammalia; Eutheria; Primates;
                      CCCGGAGGCATCCTCGCCGTCCTCAGAGTCCTGTGGCGCGGGAAGGTGCTGCAACCTCCG
                                                                                                    ccceeaaecarccrceccerccrcaeaerccrerececeeaaaecrecrecaaccrcce
                                                                                                                                                                      TTGGAGACCCTGGACGAGCTGCTGGCGGCGGCGAGCCGGCACCTTCGACGTGGCCGTG
                                                                                                                                                                                                                                          CGGCCCCTGTGGAGGCAGGCCGAGGCGAGGACCAAAGATCGACCTCCGGCTGAAGCCCGCC
                                                                                                                                                                                                                                                                                                                      CGGCCCCTGTGGAGGCAGGCCGAGGCGAGCACAAGATCGACCTCCGGCTGAAGCCCCGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quality sequence stop: 462.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="NCI_CGAP_GC6"
/note="Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; plasmid DNA from the normalized library NCI_CGAP_GC4 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 147592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="pooled germ
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="IMAGE:2476290"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="mRNA"
/db_xref="taxon:9606"
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97.9%;
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Pred. No. 2.1e-58;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 2341 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue_Procurement: Ilan Kirsch, M.D., Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; J
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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AA584408.1
                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1997)
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National Cancer Institute, Cancer Genome Anat
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                           GCCCAGCTCTTGGCCAACCTGGCGCGCTCATCCAGGCCAAGAAGGCGCTGGACCTGGGC 413
GGCCAACTTTTTGGCAAACTGGCGCGGTCATTCCAGGCCAAGAAGGCGCTGAACCTGGGC 561
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                                                                                                                                                          /clone lib="NGI_CGAP_CO9"
/note="Organ: colon; Vector: pT7T3D-Pac (Pharmacia) with a Modified polylinker; lst strand cDNA was prepared from modified polylinker; lst strand cDNA was prepared from next colon tumor, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is not normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo
                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:1090253"
                                                                                                                                                                                                                                                                                                                lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                             tissue_type="colon tumor RER+"
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                                                                               55.5%;
95.7%;
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                                                                               Score 548.6; DB 9;
Pred. No. 2.5e-58;
                                                               Mismatches
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AUTHORS
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CB194405.1
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cDNA Library Preparation: Invitrogen Corp

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)

DNA Sequencing by: Agencourt Bioscience Corporation

Clone distribution: MGC clone distribution information can

found through the I.M.A.G.E. Consortium/LLNL at:

http://image.llnl.gov

Plate: NDAM0034 row: g column: 15

High quality sequence stop: 641.
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NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
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AGENCOURT 11258413 NIH MGC 135 Mus
IMAGE:30135806 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. David Rowe
                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia; Eutheria;
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/mol_type="mRNA"
/db_xref="taxon:10090"
                                                                                                Location/Qualifiers
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Query Match
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/clone lib="NIH MGC 135"
/clone ="Wector: pCMVSport6:1; Site 1: EcoRV; Site 2: NotI;
/note="Wector: pCMVSport6:1; Site 1: EcoRV; Site 2: NotI;
/notmalized full-length enriched library from pooled mouse
embryonic limb, maxilla and mandible, day 12.5, 13.5,
14.5, and 15.5 (size selected for the 0.5-1 kb fragments)
Cloned directionally, priming method: Oligo-dT. cDNA
enrichment: >1k bp. Average insert size 1.6k bp.
Normalization (Cot value): 7.5 kb. Priming sequence:
5'GACTAGTTCTAGATCGCGAGCGCCCGCC(T)3' Tissue contributed
by, David Rowe. Library constructed by Resden, Invitrogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="IMAGE:30135806"
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Pred. No. 1.1e-57;
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Insert Length: 974 Std Error: 0.00
Seg primer: -40UP from Gibco
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Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project
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AI765658

ST2 bp mRNA linear EST 21-DEC-1999 wi82g12.xl NCI CGAP Kid12 Homo sapiens cDNA clone IMAGE:2399878 3' similar to TR:O85769 O85769 HYPOTHETICAL 24.8 KD PROTEIN. ;, mRNA
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wi82g12.x1
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1 (bases 1 to 572)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                             CAGGCCGAGGCGGAGCACAAGATCGACCTCCGGCTGAAGCCCGGCCTTGGAGACCCTGGAC
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone lib="NCI CGAP Kidl2"
/clone lib="NCI CGAP Kidl2"
/note="Organ: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Eco RI; Plasmid DNA from the normalized library NCI CGAP Kid5 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCramplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1323912-1325831, 1471368-1472903 and 1492104-1493255). Subtraction by Bento Soares and M. Fatims Boraldo. "
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/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens'
/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 1.6e-57;
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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639 bp mRNA linear EST 05-JUN-20
639 bg mRNA sequence.
67, mRNA sequence.
6910899
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Plate: LLAM10889 row: p column:
High quality sequence stop: 614.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                         /lab_host="DH10B (T1 phage-resistant)"
/clone lib="NCI_CGAP_Brn67"
/note="Organ: brain; Vector: pCMV-SPORT6; Site_1: Not1;
/note_2: Sall; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
                                                                                                                                                                                                                                                                                                      /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:4944633"
                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                     tissue_type="anaplastic oligodendroglioma"
                                                                                                                                                                                                                                                                                                                                                           organism="Homo sapiens"
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                                                                    Score 540.2; DB 12
Pred. No. 2.6e-57;
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                                                                                                                                                                                             Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geise Krizman, D., Kucaba, T., Lacy, M., Le, N., Lennon, G., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Theising, B., White, Y., Wylie, T., Waterston, R. and Washu-NCI human EST Project
Unpublished (1997)
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               IMAGE Consortium (info@image.llnl.gov)
Seq primer: -40RP from Gibco
                                                                                   Tel: 314 286 1800
Fax: 314 286 1810
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 566)
                                                   Email: est@watson.wustl.edu
This clone is available royalty-free
                                                                                                                           4444 Forest Park Parkway,
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                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Schneider fetal brain 00004" St (Stratagene); /note="Organ: brain; Vector: pBluescript SK (Stratagene); Site_1: Sst; Site_2: XhOI; Double-stranded cDNA was prepared from human fetal brain tissue. 5' and 3' adaptors were used in cloning as follows: 5' adaptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /dev_stage="5 months post-conception"
/lab_host="DH10B"
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/sex="male"
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Search completed: May 7, 2004, 19:27:15